

5 The following oligonucleotides were used in the construction of plasmids.

**Table 2: Oligonucleotides Utilized For *lovE* Variant Cloning**

<b>MO664</b> (5'GGCCATGGAGGCCGCTAGCTCGAGTCGACGGCCTAGGTGGCCAGCT3')
(SEQ ID NO:1)
<b>MO665</b> (5'GGCCACCTAGGCCGTCGACTCGAGCTAGCGGCCATGGCCGTAC3')
(SEQ ID NO:2)
<b>MO666</b> (5'GGCGCCGCTCTAGAACTAGTCTCGAGGGTACC3') (SEQ ID NO:3)
<b>MO667</b> (5'GGTACCCCTCGAGACTAGTTCTAGAGCGGCCGCC3') (SEQ ID NO:4)
<b>MO1794</b> (5'CACAGCGGCCGCTAACCTTCCCATTGGGGC3') (SEQ ID NO:5)
<b>MO1793</b> (5'CACCACTAGTACCGGGCTGATTGAC3') (SEQ ID NO:6)
<b>MO1785</b> (5'CACCACTAGTTACATTATAAAAGTAATGTG3') (SEQ ID NO:7)
<b>MO1786</b> (5'CACAGGATCCGTCATCTTGCCTCGTTATC3') (SEQ ID NO:8)
<b>MO195</b> (5'CGCGGATCCTATTGAACAAGATGGATTGCAC3') (SEQ ID NO:9)
<b>MO196</b> (5'CCGGAATTCAAGAAGAACTCGTCAAGAAG3') (SEQ ID NO:10)
<b>MO841</b> (5'ACAAAAAAAGCAGGCTCCACAATGGCTGCAGATCAAGGTAT3') (SEQ ID NO:11)
<b>MO842</b> (5'ACAAGAAAGCTGGTTCATGGAGGAATATTGTTGA3') (SEQ ID NO:12)
<b>MO2278</b> (5'GGGGATCCAATCGAGGTCCACGACCACT3') (SEQ ID NO:13)
<b>MO343</b> (5'GGGGACAAGTTGTACAAAAAAGCAGGCT3') (SEQ ID NO:14)
<b>MO2273</b> (5'GGGGATCCGCCAATGGTCCCCTCAAAC3') (SEQ ID NO:15)
<b>MO2274</b> (5'ACAAGAAAGCTGGTTCACAGAATGTTAGCTCAA3') (SEQ ID NO:16)
<b>MO344</b> (5'GGGGACCACTTGTACAAGAAAGCTGGGT3') (SEQ ID NO:17)
<b>MO2624</b> (5'GCGATGCCCAAGCGCAAGCTACGCCAATCCAGGG3') (SEQ ID NO:18)
<b>MO2654</b> (5'CGTCGCCATTGCCATTAGGCTGCGCAACTGT3') (SEQ ID NO:19)
<b>MO2680</b> (5'GGACCTTGAGCATAAATTACTATACCTCT3') (SEQ ID NO:20)
<b>MO2686</b> (5'GGCGCGTCCATTGCCATTAGGCTGCGCAACTGT3') (SEQ ID NO:21)
<b>MO2681</b> (5'TAAAACCTTGTCTTCTTCTCTAAAT3') (SEQ ID NO:22)
<b>MO2700</b> (5'CAGTGAGCGCGCGTAATACGACTCACTATAGGGCGA3') (SEQ ID NO:23)
<b>MO2701</b> (5'ATACTCTATAGACACACAAACACAAATACACACAC3') (SEQ ID NO:24)
<b>MO107</b> (5'CGCGGATCCCGTCGTTTACAAC3') (SEQ ID NO:25)
<b>MO197</b> (5'CCCAAGCTTATTATTTGACACCAGACCAA3') (SEQ ID NO:26)
<b>MO1293</b> (5'GGAAGATCTAGCATCGTGGCCAATTCTTAGTT3') (SEQ ID NO:27)
<b>MO1294</b> (5'ATAAGAATGCGGCCGCTAACCTTCCCATTGGGGCGTTGC3') (SEQ ID NO:28)
<b>MO1787</b> (5'CACAGGATCCAGCATTATTAATTAGTGTGTATT3') (SEQ ID NO:29)
<b>MO1788</b> (5'CACCACTAGTCTCGAGCAGATCCGCCAG3') (SEQ ID NO:30)
<b>MO1793</b> (5'CACCACTAGTACCGGGCTGATTGAC3') (SEQ ID NO:31)
<b>MO1794</b> (5'CACAGCGGCCGCTAACCTTCCCATTGGGGC3') (SEQ ID NO:32)
<b>MO511</b> (5'GGCCATCGATAAGTTGTACAAAAAAGCTGAAC3') (SEQ ID NO:33)
<b>MO540</b> (5'GGCGCCCTATTACACCACTTGTACAAGAAAGC3') (SEQ ID NO:34)
<b>MO1985</b> (5'CACACGTCTCCGCCCTAACCTTCCCATTGGGGCG3') (SEQ ID NO:35)

NO:35)	
<b>MO1986</b>	(5' CACACAGATCTCGTGGCCAATTCTTAGTTGA3') (SEQ ID NO:36)
<b>MO1992</b>	(5' CACACGGATCCACAATGTTACGTCTGTAGAAACCC3') (SEQ ID NO:37)
<b>MO1993</b>	(5' CACAGCGGCCGCTTCATTGTTGCCTCCCTGCTG3') (SEQ ID NO:38)
<b>MO316</b>	(5' GCGGCCGCGGCCGCCATGTCAACAAGAAT3') (SEQ ID NO:39)
<b>MO318</b>	(5' CCGCGGCCGAGTGGAGATGTGGAGT3') (SEQ ID NO:40)

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Plasmid MB2254 contains the *lovFp-HIS3p-neo* reporter gene flanked by *URA3* sequence. First primers MO664 (SEQ ID NO:1) and MO665 (SEQ ID NO:2) were annealed and 10 inserted into the *KpnI-SacI* sites of plasmid pBluescript II KS (Stratagene,). The resulting vector, MB1038, contains a *SalI* site in the polylinker. Next, the *SpeI-XhoI* fragment from pJL164 (Brachmann *et al.* *Yeast* 14:115-132 (1998)) containing a deletion of the *URA3* gene with 15 additional flanking sequences was inserted into the *NheI-SalI* sites of MB1038 to create MB1053. Primers MO666 (SEQ ID NO:3) and MO667 (SEQ ID NO:4) that contain multiple restriction sites (*NotI*, *XbaI*, *SpeI*, *XhoI* and *KpnI*) were then annealed together and ligated into the *SmaI* site of 20 MB1053 to create MB1054. Next, the following four fragments were combined in MB1054 to obtain plasmid MB2254. The *lovF* promoter from *A. terreus* genomic DNA was PCR amplified with MO1794 (SEQ ID NO:5) and MO1793 (SEQ ID NO:6) and inserted into MB1054 on a *NotI-SpeI* fragment. 25 The *HIS3* basal promoter from pRS403 (Sikorski and Hieter, *Genetics* 122:19-27 (1989)) was PCR amplified with primers MO1785 (SEQ ID NO:7) and MO1786 (SEQ ID NO:8) and inserted into MB1054 on a *SpeI-BamHI* fragment. Finally, the *neo* gene (PCR amplified with MO195 (*BamHI*) (SEQ ID NO:9) and 30 MO196 (*EcoRI*) (SEQ ID NO:10) from plasmid pYX11 (Xiao and Weaver, *Nucl. Acids Res.* 25:2985-2991 (1997)) and *CYC1* terminator sequences (*XhoI-KpnI* fragment from pRS426-GAL-S (Mumberg, *et al.*, *Nucl. Acids. Res.* 22:5767-5768 (1994))) were first combined in pRS416 (Sikorski and Hieter,

5 *Genetics* 122:19-27 (1989)) and then cut out with *Bam*HI-  
*Kpn*I and inserted into MB1054 to create MB2254.

The *lovFp-HIS3p-neo* reporter in MY2124 can confer resistance to the drug geneticin (G418). It was empirically determined that MY2124 (untransformed or  
10 transformed with parental plasmids MB2478 (*CYC1-lovE/CEN*) or MB2848 (*CYC1-lovE/At274/CEN*) was unable to grow on YPD media supplemented with 100 µg /ml G418. Plasmid MB2478 contains the *CYC1* promoter operationally linked to the entire *A. terreus* *lovE* open reading frame. The *CYC1*  
15 promoter is a relatively weak promoter and thus the *lovE* ORF in MB2478 was expressed at low levels. MB2478 was the parental vector plasmid for creating full length *lovE* variants. Plasmid MB2848 contains the *CYC1* promoter operationally linked to a chimeric open reading frame  
20 consisting of the *A. terreus* *lovE* DNA binding domain fused to the carboxy-terminal portion of the *At274* gene (U.S. Serial No. 60/257,431, filed December 22, 2000).

MB2848 was used to create *lovE* variants in which the DNA binding domain was not mutated. Both MB2478 and  
25 MB2848 contain yeast CEN and autonomously replicating sequences and both are maintained at 1-2 copies per cell. In contrast to strains transformed with MB2478 or MB2848, strains transformed with plasmid MB1644 (*TEF1-lovE/2* micron) were able to grow on G418-supplemented YPD media.  
30 The *lovE* gene of MB1644 is under control of the constitutively strong *S. cerevisiae* *TEF1* promoter. MB1644 contains a 2-micron origin for high-copy replication in yeast. An objective of these studies was to identify *lovE* variants which when expressed at low levels could confer  
35 G418 resistance similar to the highly expressed wild-type *lovE* molecule of MB1644. *S. cerevisiae* expression vectors used in these studies were constructed as follows.

MB968 is a low copy *S. cerevisiae* *URA3* based expression vector. MB968 was created by inserting the  
40 *Eco*RV fragment (containing the destination cassette) from gateway pEZC7201 (Invitrogen™, Carlsbad, CA) into